

476

SEQUENCE LISTING

<110> Aventis Research & Technologies GmbH & Co KG

<120> Spliceosomal protein and its use

<130> 199at09

<140> PCT/EP00/03949

<141> 2000-05-03

<150> DE 19925668.3

<151> 1999-06-04

<160> 18

<170> PatentIn Ver. 2.1

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<211> 18

<212> PRT

<213> Homo sapiens

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<213> Homo sapiens

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<212> PRT

<213> Homo sapiens

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Leu Ser Glu Glu
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tatttcagca ttggctctac agcgttgacag aac atg aac gat tgg atg ccc atc 234
Met Asn Asp Trp Met Pro Ile
1 5

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Ala Lys Glu Tyr Asp Pro Leu Lys Ala Gly Ser Ile Asp Gly Thr Asp
10 15 20

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Glu Asp Pro His Asp Arg Ala Val Trp Arg Ala Met Leu Ala Arg Tyr
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Val Pro Asn Lys Gly Val Ile Gly Asp Pro Leu Leu Thr Leu Phe Val
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Ala Arg Leu Asn Leu Gln Thr Lys Glu Asp Lys Leu Lys Glu Val Phe
60 65 70

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Ser Arg Tyr Gly Asp Ile Arg Arg Leu Arg Leu Val Arg Asp Leu Val
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 Thr Gly Phe Ser Lys Gly Tyr Ala Phe Ile Glu Tyr Lys Glu Glu Arg
 90 95 100

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 Ala Val Ile Lys Ala Tyr Arg Asp Ala Asp Gly Leu Val Ile Asp Gln
 105 110 115

cat gag ata ttt gtg gac tac gag ctg gaa agg act ctc aaa ggg tgg 618
 His Glu Ile Phe Val Asp Tyr Glu Leu Glu Arg Thr Leu Lys Gly Trp
 120 125 130 135

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 Ile Pro Arg Arg Leu Gly Gly Gly Leu Gly Gly Lys Lys Glu Ser Gly
 140 145 150

caa ctg aga ttt ggg gga cgg gac cgg cct ttt cga aaa cct att aac 714
 Gln Leu Arg Phe Gly Gly Arg Asp Arg Pro Phe Arg Lys Pro Ile Asn
 155 160 165

ttg cca gtt gtt aaa aac gac ctc tat aga gag gga aaa cgg gaa agg 762
 Leu Pro Val Val Lys Asn Asp Leu Tyr Arg Glu Gly Lys Arg Glu Arg
 170 175 180

cgg gag cga tct cga tcc cga gaa aga cac tgg gac tcg agg aca agg 810
 Arg Glu Arg Ser Arg Ser Arg Glu Arg His Trp Asp Ser Arg Thr Arg
 185 190 195

gat cga gac cat gac agg ggc cgg gag aag aga tgg caa gaa aga gag 858
 Asp Arg Asp His Asp Arg Gly Arg Glu Lys Arg Trp Gln Glu Arg Glu
 200 205 210 215

ccg acc agg gtg tgg ccc gac aat gac tgg gag aga gag agg gac ttc 906
 Pro Thr Arg Val Trp Pro Asp Asn Asp Trp Glu Arg Glu Arg Asp Phe
 220 225 230

aga gat gac agg atc aag ggg agg gag aag aag gaa aga ggc aag tag 954
 Arg Asp Asp Arg Ile Lys Gly Arg Glu Lys Lys Glu Arg Gly Lys
 235 240 245

aggcccaaca gcagaacccc aaagtgaagt tacagtggaa atgagtggag ggggattgtc 1014

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 Arg Ala Met Leu Ala Arg Tyr Val Pro Asn Lys Gly Val Ile Gly Asp
 35 40 45

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